

O/PE
0500
H2
1-5-01
P2

Serial Number: 09/615,039

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

RAW SEQUENCE LISTING

DATE: 08/01/2000

PATENT APPLICATION: US/09/615,039

TIME: 17:20:44

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Output Set: N:\CRF3\08012000\I615039.raw

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2   Lichtsteiner, Serge
3   Vasserot, Alain
4   Adams, Robert R.
5   Geron Corporation
7 <120> TITLE OF INVENTION: Telomerase Reverse Transcriptase Transcriptional
8   Regulatory Sequences and Methods of Using
10 <130> FILE REFERENCE: 019/251C
C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/615,039
C--> 13 <141> CURRENT FILING DATE: 2000-07-11
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/615,039

DATE: 08/01/2000

TIME: 17:20:45

Input Set : A:\Pto.amc

Output Set: N:\CRF3\08012000\I615039.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application Number
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date

OIPE

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/615,039

DATE: 07/25/2000
 TIME: 09:13:36

Input Set : A:\ES.txt
 Output Set: N:\CRF3\07252000\I615039.raw

*Does Not Comply
 Corrected Diskette Needed*

3 <110> APPLICANT: Morin, Gregg B.
 4 Lichtsteiner, Serge
 5 Vasserot, Alain
 6 Adams, Robert R.
 7 Geron Corporation
 9 <120> TITLE OF INVENTION: Telomerase Reverse Transcriptase Transcriptional
 10 Regulatory Sequences and Methods of Using
 12 <130> FILE REFERENCE: 019/251C
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 C--> 15 <141> CURRENT FILING DATE: 2000-07-11
 17 <150> PRIOR APPLICATION NUMBER: 09/244,438
 18 <151> PRIOR FILING DATE: 1999-02-04
 20 <160> NUMBER OF SEQ ID NOS: 23
 22 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

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VERIFICATION SUMMARY DATE: 07/25/2000
PATENT APPLICATION: US/09/615,039 TIME: 09:13:37

Input Set : A:\ES.txt
Output Set: N:\CRF3\07252000\I615039.raw

L:14 M:270 C: Current Application Number differs, Replaced Current Application Number
L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:726 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:428 SEQ:23
L:726 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:8
L:726 M:112 C: (48) String data converted to lower case,
L:728 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:5
L:729 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:23
L:729 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:5
L:730 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:23
L:732 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:428 SEQ:23
L:732 M:252 E: No. of Seq. differs, <211>LENGTH:Input:425 Found:428 SEQ:23